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Sequence Listing was accepted.

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Reviewer: markspencer

Timestamp: [year=2008; month=7; day=18; hr=8; min=52; sec=43; ms=165; ]

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Application No: 10591843 Version No: 1.0

Input Set:

Output Set:

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Finished: 2008-07-17 10:25:35.001  
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Total Warnings: 2  
Total Errors: 0  
No. of SeqIDs Defined: 2  
Actual SeqID Count: 2

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 402	Undefined organism found in <213> in SEQ ID (2)

## SEQUENCE LISTING

&lt;110&gt; KURUME UNIVERSITY

&lt;120&gt; Protease inhibitor and preventives or remedies for diseases

&lt;130&gt; P04AC002

&lt;140&gt; 10591843

&lt;141&gt; 2008-07-17

&lt;160&gt; 2

&lt;170&gt; PatentIn version 3.2

&lt;210&gt; 1

&lt;211&gt; 546

&lt;212&gt; DNA

&lt;213&gt; Artificial

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (70)..(540)

&lt;220&gt;

&lt;223&gt; mouse interleukin-18 and immunoglobulin

&lt;400&gt; 1

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actggtgac aac ttt ggc cga ctt cac tgt aca acc gca gta ata cgg aat 111

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5

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Ile Asn Asp Gln Val Leu Phe Val Asp Lys Arg Gln Pro Val Phe Glu

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20

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gat atg act gat att gat caa agt gcc agt gaa ccc cag acc aga ctg 207

Asp Met Thr Asp Ile Asp Gln Ser Ala Ser Glu Pro Gln Thr Arg Leu

35

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ata ata tac atg tac aaa gac agt gaa gta aga gga ctg gct gtg acc 255

Ile Ile Tyr Met Tyr Lys Asp Ser Glu Val Arg Gly Leu Ala Val Thr

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55

60

ctc tct gtg aag gat agt aaa atg tct acc ctc tcc tgt aag aac aag 303

Leu Ser Val Lys Asp Ser Lys Met Ser Thr Leu Ser Cys Lys Asn Lys

65

70

75

atc att tcc ttt gag gaa atg gat cca cct gaa aat att gat gat ata 351

Ile Ile Ser Phe Glu Glu Met Asp Pro Pro Glu Asn Ile Asp Asp Ile

80

85

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caa agt gat ctc ata ttc ttt cag aaa cgt gtt cca gga cac aac aag 399

Gln Ser Asp Leu Ile Phe Phe Gln Lys Arg Val Pro Gly His Asn Lys

95

100

105

110

atg gag ttt gaa tct tca ctg tat gaa gga cac ttt ctt gct tgc caa	447
Met Glu Phe Glu Ser Ser Leu Tyr Glu Gly His Phe Leu Ala Cys Gln	
115 120 125	

aag gaa gat gat gct ttc aaa ctc att ctg aaa aaa aag gat gaa aat	495
Lys Glu Asp Asp Ala Phe Lys Leu Ile Leu Lys Lys Lys Asp Glu Asn	
130 135 140	

ggg gat aaa tct gta atg ttc act ctc act aac tta cat caa agt	540
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Thr Asp Ile Asp Gln Ser Ala Ser Glu Pro Gln Thr Arg Leu Ile Ile
35 40 45

Tyr Met Tyr Lys Asp Ser Glu Val Arg Gly Leu Ala Val Thr Leu Ser
50 55 60

Val Lys Asp Ser Lys Met Ser Thr Leu Ser Cys Lys Asn Lys Ile Ile
65 70 75 80

Ser Phe Glu Glu Met Asp Pro Pro Glu Asn Ile Asp Asp Ile Gln Ser
85 90 95

Asp Leu Ile Phe Phe Gln Lys Arg Val Pro Gly His Asn Lys Met Glu
100 105 110

Phe Glu Ser Ser Leu Tyr Glu Gly His Phe Leu Ala Cys Gln Lys Glu
115 120 125

Asp Asp Ala Phe Lys Leu Ile Leu Lys Lys Lys Asp Glu Asn Gly Asp
130 135 140

Lys Ser Val Met Phe Thr Leu Thr Asn Leu His Gln Ser  
145 150 155